

SEQUENCE LISTING

(i) APPLICANT: Stevens, Fred J.

Argon, Yair Davis, David P. Raffen, Rosemarie

- (ii) TITLE OF INVENTION: Fibril-Blocking Peptide, A Method for Preventing Fibril Formation
- (iii) NUMBER OF SEQUENCES: 13
- (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: CHERSKOV & FLAYNIK
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 - (C) CITY: Chicago
 - (D) STATE: Illinois
 - (E) COUNTRY: United States
 - (F) ZIP: 60606
- (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: 3.50 inch, 1.4 MB storage
 - (B) COMPUTER: PC
 - (C) OPERATING SYSTEM: Microsoft Windows 98
 - (D) SOFTWARE: Wordperfect
- (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER: 09/712,819
 - (B) FILING DATE: 2000-NOV-13
- (vii) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: U.S. 60/165,424
 - (B) FILING DATE: 1999-NOV-14
- (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Cherskov, Michael J.
 - (B) REGISTRATION NUMBER: 33,664
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- (2) INFORMATION FOR SEQ ID NO: 1:
 - (i) SEQUENCE CHARACTERISTICS:

	(xi)	 (A) LENGTH: 7 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: Single (D) TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO: 1:
Phe Th	ır Leu T	Thr Ile Ser Ser 5
2)	(i)	RMATION FOR SEQ ID NO: 2: SEQUENCE CHARACTERISTICS: (A) LENGTH: 7 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: Single (D) TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO: 2:
	(xi)	
Thr Ai l	g Ile Pł	ne Ser Lys Leu 5
(2)	INFOI (i)	RMATION FOR SEQ ID NO: 3: SEQUENCE CHARACTERISTICS: (A) LENGTH: 7 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: Single (D) TOPOLOGY: linear
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 3:
Phe Ty l	yr Gln I	Leu Ala Leu Thr 5
(2)	INFOI (i) (xi)	RMATION FOR SEQ ID NO: 4: SEQUENCE CHARACTERISTICS: (A) LENGTH: 8 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: Single (D) TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO: 4:
Pro Ly 1	/s Leu I	Leu Ile Tyr Trp Ala 5
(2)	INFO	RMATION FOR SEQ ID NO: 5: (A) LENGTH: 7 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: Single

	(xi)	(D) TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO: 5:
Γhr As l	p Phe T	Thr Leu Thr Ile 5
(2)	(i)	RMATION FOR SEQ ID NO: 6: SEQUENCE CHARACTERISTICS: (A) LENGTH: 7 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: Single (D) TOPOLOGY: linear
Phe Th 1	(xi) 1r Leu I	SEQUENCE DESCRIPTION: SEQ ID NO: 6: Lys Ile Ser Arg 5
(2)	INFOI (i)	RMATION FOR SEQ ID NO: 7: SEQUENCE CHARACTERISTICS: (A) LENGTH: 9 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: Single (D) TOPOLOGY: linear
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO: 7:
Thr A	sp Phe	Thr Leu Thr Ile Ser Ser 5
(2)	(i)	RMATION FOR SEQ ID NO: 8 SEQUENCE CHARACTERISTICS: (A) LENGTH: 11 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: Single (D) TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO: 8:
Tyr G 1	ly Arg	Lys Lys Arg Arg Gln Lys Lys 5 10
(2)	INFO (i)	RMATION FOR SEQ ID NO: 9 SEQUENCE CHARACTERISTICS: (A) LENGTH: 29 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: Single (D) TOPOLOGY: linear

SEQUENCE DESCRIPTION: SEQ ID NO: 9: (xi) Tyr Glu Arg Lys Lys Arg Arg Gln Arg Arg Arg Ser Gly Ser Gly Thr Asp Phe Thr Leu Thr Ile Ser Ser Leu Gln Ala Glu Asp 25 20 INFORMATION FOR SEQ ID NO: 10 (2) SEQUENCE CHARACTERISTICS: (A) LENGTH: 29 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: Single (D) TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO: 10: (xi) Tyr Glu Arg Lys Lys Arg Arg Gln Arg Arg Arg Ser Gly Ser Gly 10 Thr Asp Ala Thr Ala Pro Ala Ser Ser Leu Gln Ala Glu Asp 25 20 INFORMATION FOR SEQ ID NO: 11: (2) SEQUENCE CHARACTERISTICS: (i) (A) LENGTH: 4 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: Single (D) TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO: 11: (xi) Ala Leu Leu Asn 1 INFORMATION FOR SEQ ID NO: 12: (1) SEQUENCE CHARACTERISTICS: (i) (A) LENGTH: 7 amino acids (B) TYPE: amino acid (C) STRANDEDNESS: Single (D) TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO: 12: (xi) Phe Thr Leu Glu Ile Ser Arg INFORMATION FOR SEQ ID NO: 13: (ii SEQUENCE CHARACTERISTICS: (ii (A) LENGTH: 7 amino acids

(B) TYPE: amino acid (C) STRANDEDNESS: Single (D) TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO: 13: (xi)

Leu Thr Leu Lys Leu Ser Arg
1 5